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(54) Title: DELIVERY AND EXPRESSION OF A HYBRID SURFACE PROTEIN ON THE SURFACE OF GRAM POSITIVE BACTERIA

(57) Abstract

Process is described for the delivery and expression of hybrid surface proteins to the surface of bacteria. The transformed bacteria are useful as vaccines, for the delivery of other active peptides to animal hosts, as diagnostic reagents and for other purposes.



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DELIVERY AND EXPRESSION OF A HYBRID SUFACE PROTEIN ON THE SURFACE OF GRAM POSITIVE BACTERIA

RELATED APPLICATIONS

This application is a continuation in part of copending application serial number 07/851,082 filed March 13, 1992 which is a continuation in part of application serial number 07/814,823 filed December 23, 1991 which is, a continuation of application serial number 07/742,199 filed August 5, 1991 which is, in turn, a continuation of application serial number 07/522,440 filed May 11, 1990. The latter two applications are now abandoned.

FIELD OF THE INVENTION

This invention relates generally to products and processes useful to deliver proteins, including protein antigens, to the surface of gram-positive bacteria and firmly attach them to the cell. More specifically, it relates to the production of a fusion protein containing at least the anchor region of the cell associated region of a gram-positive surface protein and any protein, peptide or polypeptide that may be usefully administered to an animal host. The products of the invention which comprise proteins, peptides or polypeptides, as further discussed below, placed on the surface of a

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gram positive bacteria may be used to deliver such materials to the animal host to elicit an immungenic response, for example the production of protective antibodies or for another useful purpose. protein, peptide or polypeptide may, for example, be an enzyme or other functional protein necessary for a specific purpose. It may be an antigenic determinant from a bacteria, virus, parasite or fungus. It may be a surface antigen from a mammalian tumor cell or of male sperm. It may be an allergen such as a vespid venom. The invention also relates to novel plasmids, genes, chromosomes and transformed bacteria employed in the production of such fusion proteins. The invention, additionally, provides novel vaccines which employ gram-positive bacteria designed to deliver to an animal host a foreign antigen normally present on a pathogenic microorganism which is associated with the virulence of that pathogen and which will elicit antigens to protect the host against infection or disease caused by the pathogenic microorganism.

The products of the invention are also useful as diagnostic agents.

- 3 -

The invention also provides a means to deliver enzymes, placed on the bacterial surface, to specific areas of interest.

The term "animal" as used herein refers to living beings including mammals such as man; bovines, especially beef cattle; sheep and goats; poultry, especially chickens, ducks and turkeys; as well as fish, especially those raised in fish farms such as salmon, trout and catfish. This invention is of special importance to mammals.

BACKGROUND OF THE INVENTION

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The essence of this invention is that it provides a method for the production of novel non-pathogenic gram positive bacteria expressing a hybrid surface protein which may be a hybrid surface antigen, comprising two principal parts, an anchor segment comprised of amino acid residues and an N-terminal active polypeptide segment, both of which will be defined and discussed in more detail below.

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This invention will be better understood by consideration of the M protein and its structure.

The M protein is a coiled coil surface antigen which is the virulence factor of group A streptococci, a gram positive bacteria. The M protein of Streptococcus pyogenes of M type 6 contains 441 amino acid residues.

Its structure will be discussed in more detail below, but it will be useful to discuss the cell associated region at this point. Using the standard one letter representation of amino acids, the structure of the anchor region of the cell associated region of the M6 protein from amino acid residue 407 to residue 441 may be represented as:

15 LPSTGETANPFFTAAALTVMATAGVAAVVKRKEEN

Reading from the first leucine residue (L) at the N-terminal of the anchor region, the region includes the LPSTGE segment; a spacer segment containing three amino acid residues TAN; a hydrophobic segment of twenty amino acids, PFFTAAALTVMATAGVAAVV; followed by a highly charged tail segment, KRKEEN.

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It has been observed as a result of structural studies of a large number of surface proteins of gram positive bacteria that the above described anchor region of the M6 protein is highly conserved among all known surface proteins of gram positive bacteria. Many of them are shown by Fischetti et al (Reference 1). Generally, the hydrophobic segment contains about 15 to 20 amino acid residues, the charged tail segment about 4 to 6 amino acid residues, and the spacer segment from about 3 to 6 amino acid residues. Most remarkable however, is the high degree of homology, practically 100% in the LPSTGE segment of the known surface proteins. The variations that occur are almost exclusively at the 3 and 6 positions. Therefore, the region may be generally represented as LPXTGX.

The following Table 1 shows the remarkable extent of this homology thus far established amongst forty different surface proteins of gram positive bacteria.

- 6 -

TABLE 1
SEQUENCED SURFACE PROTEINS FROM GRAM-POSITIVE BACTERIA

				•	
	NAME/GENE	SURFACE PROTEIN	ORGANISM	LPSTGE	<u>REF</u>
	1. M6	M protein	S. pyogenes	LPSTGE	(2)
5 ′	2. M5	M protein	S. pyogenes	LPSTGE	(3)
	3. M12	M protein	S. pyogenes	LPSTGE	(4)
	4. M24	M protein	S. pyogenes	LPSTGE	(5)
	5. M49	M protein	S. pyogenes	LPSTGE	(6)
	6. M57	M protein	S. pyogenes	LPSTGE	(7)
10	7. M2	M protein	S. pyogenes	LPSTGE	(8)
	8. ARP2	IgA binding protein	S. pyogenes	LPSTGE ·	(8)
	9. ARP4	IgA binding protein	S. pyogenes	LPSTGE	(9)
•	10. FCRA	Fc binding protein	S. pyogenes	LPSTGE	(10)
	11. Prot H	Human IgG Fc binding	S. pyogenes	LPSTGE	(11)
15	12. SCP	C5a peptidase	S. pyogenes	LPTIND	(12)
	13. T6	Protease resistant	S. pyogenes	LPSTGS	(13)
		protein			
	14. bac	IgA binding protein	Gr. B strep	LPYTGV	(14)
	15. Prot G	IgG binding protein	Gr. G strep	LPTTGE	(15)
	·				
20	16. PAc	Surface protein	S. mutans	LPNTGE	(16)
	17. spaP	Surface protein	S. mutans	LPNTGE	(17)
	18. spaA	Surface protein	S. sobrinus	LPATGD	(18)
	19. wapA	Wall-associated	S. mutans	LPSTGE	(19)
		protein A		•	•
25	20. Sec10	Surface protein	E. fecalis	LPQTGE	(20)
	21. Asc10	Surface protein	E. fecalis	LPKTGE	(20)
	22. asa1	Aggregation substance	E. fecalis	LPQTGE	(21)

	23.	Prot A	IgG binding protein	s.	aureus	LPETGV	(22)
si.	24.	FnBP	Fibronectin binding protein	s.	aureus	LPETGG	(23)
	25.	wg2	Cell wall protease	s.	cremoris	LPKTGE	(24)
5	26.	In1A	Internalization	L.	monocyto-	LPTTGE	(25)
			protein	gei	nes		
	27.	Fimbriae	Type 1 fimbriae	A.	viscosis	LPLTGA	(26)
	28.	Fimbriae	Type 2 fimbriae	A.	naeslundii	LPLTGA	(27)
	29.	Mrp4	IgG/Fibrinogen binding	s.	pyogenes	LPSTGE	(10)
10	30.	sof22	Serum opacity factor	s.	pyogenes	LPASGD	(54)
	31.	Sfb	Fibronectin binding	s.	pyogenes	LPATGD ·	(55)
	32.	Prot L	Light chain binding	P.	magnus	LPKAGS	(56)
	33.	bca	alpha antigen	Gr	. B strep	LPATGE	(57)
	34.	fnbA	Fibronectin binding	s.	dysgalactiae	LPQTGT	(58)
15	35.	fnbB	Fibronectin binding	s.	dysgalactiae	LPAAGE	(59)
	36.	EmmG1	M protein	Gr.	. G Strep	LPSTGE	(60)
	37.	DG12	Albumin binding				
			protein	Gr.	. G strep	LPSTGE	(61)
	38.	MRP	Surface protein	s.	suis	LPNTGE	(62)
20	39.	FnBp	Fibronectin binding				
			protein	s.	aureus	LPETGG	(63)
	40.	cna	Collagen binding				F
			protein	s.	aureus	LPKTGM	(64)

It is apparent that this highly homologous region of the

surface proteins of gram-positive bacteria is essential to
anchoring bacterial surface proteins to the cell (28). This
segment, which is referred to herein as the LPXTGX segment, is the
crucial segment of the cell associated region of surface protein
for anchoring the proteins to the surface of gram positive

bacteria.

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This discovery has been confirmed by Schneewind et al (65).

Using the Protein A molecule of <u>Staphlococcus aureus</u>, these investigators have established that the complete complex (LPXTGX motif, hydrophobic domain and charged tail) are necessary to deliver the Protein A molecule to the cell surface and that changes in the LPXTGX motif or deletion thereof will not inhibit expression of the molecule but will prevent it from anchoring to the surface.

BRIEF DESCRIPTION OF THE DRAWINGS

Fig. 1 shows the alignment of the amino acids in the C-terminal regions of a variety of surface antigens from gram-positive bacteria and the gene segments which are employed to express them. The LPSTGE motif is shaded and the proteins were aligned along this consensus sequence. In 10 out of 11 proteins, a conserved lysine residue was found 2 or 3 residues preceding the consensus LPSTGE sequence (boxed). The homologous carboxyterminal hydrophobic regions are also boxed. Abbreviations used in the left column are the same as those in Table 1.

Fig. 2 a model of the M protein.

Fig. 3 shows the complete amino acid sequence of the M6 protein.

Fig. 4. shows the host vector system GP232-pVMB20.

PCT/US93/02355

WO 93/18163

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Fig. 5 shows the construction and chromosomal integration of the M6:E7 translation fusion in GP246.

Fig. 6 shows the results of studies designed to demonstrate that the M6:E7 fusion protein expressed in Streptococcus gordonii GP246 located on its surface.

Fig. 7 shows that cell extracts of recombinant Streptococcus gordonii 246 expresses the M6:E7 fusion protein.

Fig. 8 shows that animals immunized with the recombinant GP246 produce antibodies reactive with the E7 protein.

Fig. 9 shows a universal plasmid of the invention.

Fig. 10 shows a typical hybrid protein of the invention.

Figs. 11 and 12 show the results of studies in which mice were treated with a hybrid protein of the invention.

Careful study of Figs 2 and 3 will assist in understanding this invention. Fig. 2 represents a model of the M protein with certain of the positions and segments identified. Fig. 3 identifies each amino acid residue in the complete structure of the M6 protein.

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It will be understood from the foregoing discussion that the segment of the M protein identified as the cell associated region in Fig. 2 is analogous to regions for surface proteins found on other gram-positive bacteria such as those listed in Table 1. As discussed above, there is a high degree of homology particularly within the anchor sequence especially the LPXTGX segment found in all grampositive surface proteins. As will be described below, and for the purpose of illustrating this invention, the gene segment spanning region 122 to 300 of the gene which expresses the M protein is genetically removed and replaced by a new gene segment expressing a foreign protein, for example, an antigen which will generate useful antibodies, thereby producing a novel hybrid surface protein. Because of the great degree of homology within the cell associated regions of all gram-positive bacteria, the hybrid gene can be employed in any gram positive bacteria. The selected bacteria will express the designed hybrid protein using the anchor region to attach the molecule to the cell and positioning the inserted active segment on the cell surface. The inserted active segment is hereinafter referred to as the "active polypeptide".

- 11 -

The term "active polypeptide" is used herein in the broadest possible sense. It refers to any peptide, polypeptide or protein which may be delivered to an animal host for any useful purpose. For example, as described in detail hereinafter, the cell associated region of a protein from gram-positive bacteria can be fused to a segment of a viral protein from a pathogen to produce a hybrid surface protein which will be expressed by non-pathogenic bacteria. The bacteria can colonize an animal host and function as a vaccine. It will elicit antibodies in an animal host to protect against or inhibit subsequent infection by the virus. The fused viral segment is the "active polypeptide" of that particular embodiment of the invention.

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For convenience, the term "polypeptide" will hereinafter be used to refer to molecules which are of sufficiently high molecular weight to be called proteins, to products of lesser molecular weight, usually called polypeptides, and to products of even lesser molecular weights normally referred to as peptides.

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The "active polypeptide" may be any polypeptide which can be delivered to an animal host for a useful purpose. It could be, for example the viral segment referred to above. It could also be an antigen, from any pathogenic virus, or from a bacteria, parasite or fungi. In such instances, the active polypeptide may be the complete antigen, the antigenic determinant of the antigen or a segment of the antigen which includes the antigenic determinant. The useful purpose will be to elicit a protective immune response by the production of antibodies or to elicit a cellular immune response to the antigen.

The term "cell associated region" as used in this specification and claims will be readily understood by reference to the figures, especially Figs. 2, 3 and 10. It will be seen that the cell associated region includes a cell wall spanning sequence of amino acid residues and a carbohydrate spanning segment. The wall spanning sequence may be omitted from the hybrid proteins of this invention, although it is presently preferred not to do so.

The cell associated region also includes the anchor region which contains the anchor segment (LPXTGX), the spacer segment, the hydrophobic segment and the charged tail segment. The anchor sequence is essential to the hybrid proteins of this invention. Without the, sequence the hybrid protein will not be retained on the surface of the gram positive bacterial carrier.

Fig. 10 shows the balance of a typical hybrid

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protein of the invention which, as shown, includes the active polypeptide fused to the cell associated region. The figure also shows a leader segment and a small N-terminal segment at the amino terminal of the active polypeptide. These segments cooperate in the proper placement of the hybrid protein on the surface of the gram positive bacteria. The function of the leader segment is to permit the proper processing of the hybrid protein within the cell. The N-terminal segment in cooperation with the leader segment enables the leader peptidase enzyme to separate the leader segment from the N-terminal segment and permit the hybrid protein to assume its proper position on the surface of the bacteria. The leader segment and the N-terminal segment are from the same surface protein

as the protein of the cell associated region. The N-terminal segment suitably contains the first few amino acid residues derived from the amino end of the protein used in the cell associated region of the hybrid protein. About ten such residues are normally sufficient for the proper processing, but segments containing up to twenty or more amino acid residues may be employed.

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The active polypeptide is not necessarily an antigen from a pathogenic organism. It may also be an enzyme, for example. In that event, the useful purpose will be to deliver the enzyme on the surface of a non-pathogenic bacteria to a specific site in the animal host colonized by that non-pathogenic bacteria.

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The active polypeptide may be the complete molecule which comprises the enzyme. Alternatively, it may be only that segment of the polypeptide which is required to accomplish the useful purpose.

- 15 -

The active polypeptide may be an antigen which will react with antibodies elicited by pathogenic microorganisms. Bacteria of the invention carrying such active polypeptide antigens are useful as diagnostic reagents.

When reference is made herein to non-pathogenic bacteria it should be understood to include gram positive commensal bacteria as well as pathogenic bacteria which have been modified or attenuated by the usual procedures so that their pathogenicity is weakened or destroyed and which express hybrid proteins in accordance with the process described herein, such hybrid proteins containing a C-terminal region including the LPXTGX region.

BRIEF DESCRIPTION OF THE INVENTION

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This invention provides non-pathogenic gram positive bacteria which express at least one hybrid surface protein the C-terminal region of which is attached to the bacteria. The C-terminal region includes, but is not limited to, the cell associated region of a surface protein normally expressed by a gram-positive bacteria which may be pathogenic or non-pathogenic. The balance of the hybrid surface

- 16 -

protein includes an active polypeptide which may be delivered to an animal host for any useful purpose. The surface protein "normally produced" by the bacteria of this invention refers to the whole protein produced by the bacteria before it is genetically altered in accordance with the process of this invention.

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As is known from previous studies (29) and indicated in Figs 2 and 3, amino acids 298-441 of the M protein are buried within the cell whereas residues 1-297 are exposed on the surface of the bacterial cell. In accordance with this invention, by employing genetic manipulations described and illustrated herein, nearly all of the surface exposed region of the M protein may be removed and replaced by an active polypeptide that is linked to the C-terminal cellassociated region of the M protein to produce a hybrid surface protein of the invention. Because the cell associated region is essentially the same for all gram-positive surface molecules (Fig. 10), a smimilar strategy may be used whereby the cell associated region of any one of these molecules may be used as a delivery carrier or vehicle to deliver an active polypeptide from another source. The new gene

- 17 -

employed to express the hybrid surface protein may be inserted into a plasmid by standard techniques and the plasmid used to transform an Escherichia coli or other vector. The E. coli thereafter will express the new fusion protein, i.e. the hybrid protein of this invention. The E. coli procedure is useful for the production of large amounts of hybrid protein which can be isolated from the periplasm of this gram negative organism. However, for most utilities of this invention, it is preferred to transform a gram positive organism for the production of the hybrid protein on the surface of the gram positive bacteria. The recombinant genes produced in accordance with the invention may be processed by any gram positive bacteria.

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Alternatively, the newly constructed plasmid containing the fusion gene may be used to integrate the fusion gene into the chromosome of a gram-positive bacteria, for example the chromosome of Streptococcus mutans. The newly produced strain of Streptococcus mutans will thereafter produce the new hybrid protein of this invention by expressing it on the cell surface. This is the presently preferred procedure for producing the products of this invention.

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Antigenic polypeptides from a wide variety of microorganisms may be employed as the active polypeptides of the invention. These include pathogenic microorganisms which infect man and animals. There follows a representative list of typical microorganisms which express polypeptides useful in the practice of this inventions. The transformed bacteria of the invention may be used to treat or prevent the diseases associated with infection by the microorganism.

Fungi: Candida albicans, Aspergillus fumigatus,

Histoplasma capsulatum (all cause disseminating

disease), microsporum canis (animal ringworm)

Parasitic Protozoa: <u>Plasmodium falciparum</u>

(malaria), <u>Trypanosoma cruzi</u> (Sleeping sickness),

Spirochetes: <u>Borrelia bergdorferi</u> (Lyme disease), <u>Treponema pallidum</u> (syphilis), <u>Borrelia recurrentis</u> (recurring fever), <u>Leptospira icterohaemorrhagiae</u> (leptospirosis)

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Bacteria: Neisseria gonorrhoeae (gonorrhoea),

Staphylococcus aureus (endocarditis),

Streptococcus pyogenes (rheumatic fever),

Salmonella typhosa (salmonellosis), Hemophilus

influenzae (influenza), Bordetella pertussis

(whooping cough), Actinomyces israelii

(actimomyosis), Streptococcus mutans (dental caries)

Streptococcus equi - Strangles (Horses),

Streptococcus agalactiae (bovine mastitis),

Streptococcus anginosus (canine genital infections)

Viruses: Human immunodeficiency virus (HIV),
Polio virus, Influenza virus, Rabies virus,
Herpes virus, Foot and Mouth Disease virus,
Psittacosis virus, Paramyxovirus, Myxovirus,
Coronovirus

In one embodiment of this invention, a nonpathogenic gram-positive bacteria that is a commensal
organism for the host animal will be used to express
the hybrid protein on its surface, by inserting the
gene coding for the hybrid protein into the non-

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pathogenic gram-positive bacteria. If the fusion is with an active polypeptide that is an antigen from a pathogenic organism (bacterial, viral, parasitic, or fungal), and the resulting commensal bacteria expressing the hybrid antigen is administered to an animal host, there will be an immunological response by both humoral and cell mediated pathways. One possible immunological response is the production of antibodies, thereby effecting protection against infection by the pathogen. If the fusion is with an enzyme, the enzyme will be expressed on the surface of the commensal. A wide variety of active polypeptides may be delivered to the surface of gram positive bacteria as will be apparent to the skilled artisan from a study of this disclosure.

The genes, plasmids, chromosomes, and transformed bacteria of this invention are produced by the application of standard recombinant techniques well known to those skilled in the art. For example, an oligonucleotide or a restriction fragment coding for a viral coat protein, the virulence factor of a bacterial surface antigen (or, in fact, the complete antigen), an enzyme or other desired progenitor of a hybrid surface protein of the invention may be ligated

to the gene which expresses the cell associated region of the M protein or other known surface protein from a gram positive bacteria such as those listed in Table

1. The resulting hybrid gene will be used to transform a gram positive bacteria to effect the production of a surface hybrid protein of the invention.

This invention will be further explained in conection with the expression of the hybrid protein M6:E7 by the GP246 strain of Streptococcus gordonii. This hybrid antigen contains the C-terminal cell associated segment of the M6 protein. The antibody eliciting polypeptide, E7, is an early protein from a papillomavirus strain.

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Streptococcus gordonii is a commensal bacteria of the oral cavity. E7 is a 98 amino acid early protein produced during viral replication of human papillomavirus type 16 (HPV16). It is an oncoprotein, antibodies to which are found in patients with cervical cancer. It is considered a major candidate antigen for vaccines against HPV-induced malignant neoplasias.

- 22 -

The novel bacteria of this invention may be produced by standard homologous recombination shown in Figs. 4 and 5. As illustrated, a new strain of Streptococcus gordonii, GP232 which expresses large quantitites of M6 protein was employed as the host to produce a new strain GP246.

The procedure employed as illustrated in Figs. 4 and 5 is standard and widely utilized by the skilled practitioner. The technology, as illustrated, is to construct a gram positive recipient organism for the insertion of the fusion gene in a random location in the chromosome that results in a high expression of the fusion gene. The insertion site may vary from organism to organism and even within the same organism. The advantage of the method is that it insures that the inserted gene is in a site which will result in high expression of the fusion gene, rather than rely on the expression as a result of a foreign promoter. Thus, the method assures that a region with a strong promoter is selected.

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In the process of producing the novel GP246 strain, the 538 base pair segment between sites KpnI and HindIII of the M6 protein was excised from the

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novel plasmid pVMB20, produced as described below, and the 294 base pair encoding for E7 was inserted in its place to produce the novel gene which was utilized for the production of the new plasmid pVMB21. The region of the M protein that was excised results in the placement of the E7 molecule at the cell surface with a 120 amino acid segment of the M6 N-terminus including the leader segment fused to the N-terminus of E7. The construct was made in E. coli and the recombinant plasmid was used to transform S. gordonii by the standard double cross-over procedure with GP232 to effect chromosomal integration of the gene which expresses the M6:E7 protein by the new strain GP246.

MATERIALS AND METHODS

Recombinant DNA techniques. Gene fusions in

Escherichia coli vectors were obtained and constructed by standard procedures (30).

Streptococcal transformation. Frozen cells of naturally competent <u>S. gordonii</u> Challis were prepared and transformed by known procedures (31). Plating and

scoring of transformants on multilayered plates was performed using erythromycin at a concentration 5 ug/ml in the overlay using known procedures (32).

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Genetic analysis of transformants. Transformants were streaked on the surface of 3 blood agar plates by toothpick transfer of colonies from the selection plate. The first plate contained erythromycin (5 ug/ml), the second chloramphenicol (5 ug/ml), and the third had no antibiotic. Plates were incubated 36 hours at 37°C. After incubation, a nitrocellulose membrane was applied to the surface of the plate that contained no antibiotic, and kept for 20 min. at room temperature. The membrane was then incubated 30 min at 37°C and 15 min at 80°C in a vacuum oven. The presence of E7 protein bound to the nitrocellulose was detected using anti-E7 polyclonal antibodies (1:5000 dilution), obtained from rabbits immunized with a MS2:E7 fusion protein produced in E. coli (33).

Immunofluorescence. Bacteria grown in ToddHewitt broth (Difco) were harvested in late
exponential phase, applied on a glass slide, and fixed
with methanol. Slides were treated with M6- or E7specific antibodies (1:50 dilution), extensively

- 25 -

washed, and then reacted with goat anti-rabbit (or anti-mouse) IgG antiserum (1:100 dilution), conjugated with rhodamine. Results were observed under a Confocal Fluorescence Imaging System MRC-500 Bio-Rad microscope.

Western blot analysis of cell extracts.

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Streptococci were grown to late stationary phase in Todd-Hewitt borth (Difco). Cells were harvested and resuspended in 50 mM Tris(pH 8.0), 50 mM MgCl₂, 30% sucrose. Protoplasts were obtained by treating the cell suspension with lysozyme (100 ug/ml) for 30 min. at 0°C. Protoplasts were then centrifuged and resuspended in 50 mM Tris (pH 8.0). Thorough lysis was achieved by five cycles of quick freezing/thawing of the suspension. Cells that did not lyse and gross debris were discarded by low speed centrifugation at 1,000 rpm for 15 min, whereas the supernatant, containing membranes and cytoplasm, was used for Western blot analysis. The extract obtained from about 5x10⁸ streptococcal cells was run on a gel, and Western blot was performed by convenitonal methods.

- 26 -

Immunization of mice. Balb/c mice were immunized subcutaneously with 5X10⁸ live GP246 streptococcal cells (5X10⁷ colony forming units) emulsified in complete Freund's adjuvant. Two and three weeks after the primary immunization, animals were given subcutaneous boosters of the same bacterial dose emulsified in incomplete Freund's adjuvant. Animals were bled a week after the last boost.

RESULTS

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Production of the plasmid pVMB3: A construct was made where ermC, a gene conferring resistance to erythromycin (34), was cloned adjacent to emm-6.1, so that both genes would be within the same ClaI fragment. In this construct the initiation codon of emm-6.1 was 19 bp downstream of one of the ClaI sites, so that ClaI cleavage would leave emm-6.1 promoterless. Plasmid pVV3:M6 (35), containing this ClaI site upstream of the emm-6.1 coding sequence, was used in the experiments. The 2.0 kb MspI fragment of pE194, containing ermC, was ligated with a partial ClaI digestion of pVV3:M6. After transformation of E. coli DH5, a clone was isolated with a plasmid, pVMB3, containing the ClaI fragment.

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Production of the strain GP230: The 3.4-kb ClaI fragment of pVMB3, containing ermC and the promoterless emm6.1, was ligated with chromosomal DNA of S. gordonii also cut with ClaI. The ligation mixture was used to transform the naturally transformable S. gordonii "Challis", strain V288. this method the emm-6.1/ermC ClaI fragment was integrated at random into the chromosome. The chromosomal DNA ligated to the ClaI fragment provided the homology for integration during transformation. Erythromycin-resistant (Em-r) transformants were selected and analyzed for production of M6 protein by "streak blot". Of 700 Em-r transformants, 196 (28%) produced M6 protein. Based on the semiquantitative dot blot analysis, GP230 appeared to be the best M6 producer.

Production of the strain GP232: S. gordonii

GP230 was transformed with the chromosomal DNA of

pneumonococcal strain GP69. This pneumonoccal strain

is resistant to chloramphenicol but susceptible to

erythromycin produced according to the procedure of

Pozzi and Guild (36). When GP69 chromosomal DNA is

used to transform GP230, recombination occurs at the

level of the ermC sequence leading to insertion of the

- 28 -

CAT sequence into the copy of ermC integrated into the GP230 chromosome. This insertion yields GP232 which, as discussed and shown in the figures, expresses M6 on its surface and is also resistant to chloramphenical and sensitive to erythromycin.

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Expression of E7 protein of HPV16 in <u>S. gordonii</u>:
The integration vector, pVMB20, was constructed to allow insertion of heterologous DNA sequences into the <u>emm-6.1</u> gene present on the chromosome of GP232.
pVMB20 is a novel <u>Escherichia coli</u> plasmid that does not replicate in <u>Streptococcus</u> and carries emm-6.1 and the erythromycin resistance marker ermC. The host-vector system GP232-pVMB20 is shown in detail in Figs. 4 and 5. Specifically, the novel plasmid pVMB20 and novel bacteria <u>S. gordonii</u> GP246 were prepared by the following procedure.

expression: [A] In the chromosome of the novel host strain <u>S. gordonii</u> GP232, a copy of the M6 protein gene (<u>emm-6.1</u>)(37), promoterless but with its own ribosomal binding site was integrated downstream of a strong chromosomal promoter. Adjacent to <u>emm-6.1</u> is found <u>ermC</u> (34), whose coding sequence is interrupted

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by insertion in its BcII site of the 1.8-kb MboI fragment of pC221 containing a cat gene (38). GP232 expresses M6 on its surface and is resistant to chloramphenical and sensitive to erythromycin. obtained using transformation to integrate heterologous DNA into the streptococcal chromosome. The structure is shown in the figure. The size (5.2kb) of the heterologous DNA integrated into the chromosome in GP232 was determined by Southern blot analysis. The integration vector pVMB20, is a 6.3-kb E. coli plasmid that does not replicate in Streptococcus. It was obtained by subcloning in pBLUESCRIPT (Stratagene, La Jolla, California) a 3.4kb ClaI fragment of plasmid pVMB3 containing emm-6.1 and ermC by the procedure explained below. This is the same ClaI fragment which is integrated into the chromosome of GP232, the only difference being that in GP232 ermC is interrupted by cat. When pVMB20 is used as donor DNA in transformation of competent cells of S. gordonii GP232, erythromycin-resistant transformants are obtained by recombination between the integration vector and the homologous chromosomal sequences. The DNA fragment containing the cat gene is deleted in the chromosome of these transformants, whereas an intact ermC gene is restored.

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(B) The E7 protein gene of HPV16(39) was cloned into the emm-6.1 sequence of pVMB20 to yield pVMB21. pVMB20 was digested with KpnI and HindIII and ligated with a KpnI/HindIII segment containing the E7 sequences obtained by in vitro DNA amplification (polymerase chain reaction) performed on plasmid pMBS21L/E7 (33). Amplification primers were designed in order to obtain "in frame" insertion of the 294 bp encoding for E7 into emm-6.1. Nucleotide sequence analysis of pVMB21 confirmed the expected structure of the M6:E7 translational fusion. pVMB21 was linearized and used to transform GP232. E7 was found to be expressed in 6% of the erythromycin-resistant transformants. In these transformants integration of the pVMB21 sequences produced a deletion involving the cat gene. The structure of GP246, a representative transformant, was confirmed by Southern blot analysis. The nucleotide sequence of the junction fragments of the M6:E7 gene fusion present on the chromosome of GP246 was also determined after cloning in pBLUESCRIPT the ClaI fragment containing the M6:E7 fusion.

Surface expression of the E7 protein: Expression of the E7 protein of HPV16 in <u>S. gordonii</u> on the surface of strain GP246 was verified by

- 31 -

immunofluorescence using antibodies specific for either the M6 protein carrier or the E7 insert.

GP246, containing the M6:E7 gene fusion exhibited positive fluorescence when reacted with either M6-specific or E7-specific polyclonal antibodies (Fig. 6), confirming the surface location of the E7 molecule and the M protein on the surface of S. gordonii. No fluorescence was observed when GP246 was reacted with known monoclonal antibody 10A11, which is specific for an epitope of M6 whose coding region was contained in the KpnI/HindIII fragment deleted in the construction of M6:E7 gene fusion (Fig. 6).

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To demonstrate that the E7-expressing recombinant streptococci in fact produce an M6:E7 fusion protein, S. gordonii cell extracts were analyzed by Western blot (Fig. 7). In cell extracts of GP246, the same bands reacted with E7- and M6-specific antibodies, whereas no E7-specific reactivity was found in the recipient GP232, whose extracts showed M6-specific reactivity (Fig. 7).

Substantially the same procedures used to construct M6:E7 a hybrid protein containing the C-terminal region of the M6 molecule and allogen 5 was

- 32 -

successfully expressed on the surface of <u>S. gordonii</u>. Western blots of the cell wall extract using allergen-5 specific antibodies established (as with M6:E7) that allergen-5 was indeed expressed in the cell wall fraction of the gordonii. Allergen 5 was obtained as described by Fang, et al (66).

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Immune response to fusion protein on streptococcal surface: The immunogenicity of the M6:E7 fusion protein was examined by immunizing mice with recombinant S. gordonii GP246 expressing the M6:E7 fusion protein. Control mice were immunized with the isogenic strain GP232 which express M6 protein. Sera from three animals immunized with each strain were pooled and tested by Western blot for their reactivity with purified E7 protein produced in Schizosaccharomyces pombe. Fig. 8 shows that animals immunized with strain GP246 containing surface M6:E7 produced antibodies reactive with the E7 protein, indicating that E7 protein is immunogenic when expressed as a fusion protein on the streptococcal surface. No antibodies to the E7 protein were seen in the sera of mice immunized with GP232 containing only M6.

- 33 `-

The procedure which has been described permits the production of the non-pathogenic bacteria S. gordonii GP246 or other transformed gram positive bacteria which express a hybrid surface protein such as the novel hybrid surface antigen M6:E7. This novel antigen is a hybrid protein, the carboxy terminus of which is attached to the bacteria and is substantially the same C-terminal region normally found in other gram-positive surface proteins as explained above. The active polypeptide of this hybrid surface protein is the antigen E7 of HPV 16. When the bacteria is administered, e.g. by colonization, to an animal host in need of protection, the active polypeptide will elicit both humoral and cell mediated responses resulting in the production of a protective immune response against infection by papillomavirus.

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It will be noted that the heterologous antigen of the hybrid surface protein of this invention prepared as described above is a viral antigen produced during viral replication. Thus, the process of this invention makes possible the production of novel gram positive bacteria for the delivery of viral antigens

to an animal in sufficient quantities to elicit a protective immunogenic response to infection by a virus.

The novel M6:E7 and novel intermediate products of this invention are produced from two starting materials. These are pVV3:M6 and GP69. They can be prepared by the procedures described in the cited references. Therefore the novel products can all be obtained from known materials utilizing the procedures described herein. However, to assist in the practice of the invention and without admitting any necessity to do so pVV3:M6 and GP69 have been deposited at the American Type Culture Collection under the accession numbers ATCC 68003 and ATCC ______, respectively.

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The foregoing M6:E7 example illustrates the utilization of a non-pathogenic commensal gram positive bacteria normally present in the mammalian oral cavity to protect the whole body against viral infection. Analogously prepared non-pathogenic bacteria can be used to express and deliver other useful antigens by the processes described are illustrated herein. For example, protein antigens on the surface of mammalian tumors can be fused with the

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cell wall associated region of a surface antigen of any gram positive bacteria, the fusion gene inserted in a gram positive commensal, and the resulting bacteria employed as a vaccine to generate tumor specific immune responses useful in tumor therapy.

These examples illustrates only of one aspect of the invention. The invention, in fact, provides a delivery system for any kind of polypeptide which may be useful to raise an immune response in an animal or for other useful purposes.

When utilized as a vaccine, the selected transformed commensal will be employed to colonize a mammal. It will produce the selected hybrid surface protein, the active polypeptide segment of which will elicit the production of protective antibodies.

Gram positive non-pathogenic bacteria normally found on the vaginal mucosa may be employed as contraceptives. For this utility, surface antigens of male sperm can be employed as the active polypeptide on the hybrid surface protein of the bacteria

Lactobacillus acidophilus normally found on the surface of the vaginal or uterine mucosa. When the

transformed bacteria are utilized to colonize such mucosa, the hybrid proteins will elicit the production of antibodies to the surface antigens of the sperm and bring about inactivation of the sperm. In effect, the sperm will be viewed by the female body as a foreign substance and the preformed antibodies will react with and inactivate the sperm cells.

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There are two important advantages to this type of contraception. One is that, since the sperm will be inactivated, the ova will not be fertilized. The other is that the contraceptive effect can be neutralized simply by treating the immunized female with an antibiotic to clear the gram positive bacteria expressing the hybrid antigen.

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Still another utility of the transformed bacteria of this invention is the delivery of a surface antigen from an HIV virus to the surface of a gram-positive commensal organism to provide a vaccine which may be used to prevent AIDS. For this utility, the polypeptide GP120, the V3 loop of GP120, GP160 or a related surface antigen or segments of such antigens such as sequence 735 to 752 of GP16 or RP135 a 24-amino acid segment of GP 120 from the HIV virus.

- 37 -

These polypeptides will be fused to the leader and at least a portion of the N-terminal sequence of a gram positive surface protein and its anchor region for delivery to the surface of the commensal bacteria. When the transformed commensal expressing the HIV antigens on its surface is delivered to a susceptible mammal, an immune response is raised to protect against infection by HIV virus. Furthermore, by using a recombinant vaginal commensal to deliver an HIV antigen such as GP120 to the vagina, the IgA antibodies produced to this antigen are protective against infection at this site. IgA antibodies in vaginal secretions will reduce the number of HIV particles in the secretions of an HIV positive female and thus reduce the spread of AIDS.

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The products of this invention are useful in densitization therapy. This type of therapy is widely employed to alleviate the discomfort of atopic individuals who develop exaggerated IgE responses to antigens (called allergens in the allergy field). The increased production of IgE antibodies is believed to be a principal cause of the allergic responses such as hay fever, asthma, and anaphylactic shock.

The allergens may arise from any of a variety of sources including foods, molds, dust and animal furs. Flora such as roses or rag weed are a major source of allergens. The "sting" of vespids such as wasps, yellow jackets and hornets contain allergenic proteins.

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The conventional therapy employed to ameliorate the immune response from exposure to allergens has been hyposensitization treatment which involves repeated injections of increased doses of allergen. This causes an increase in allergen specific IgG antibodies which blocks the binding of allergen specific IgE to mast cells.

A number of allergens-specific peptides and proteins have been identified, isolated, and cloned. Those include, for example, allergin 5, a vespid venom protein. These allergens can be included as the active polypeptide of the hybrid surface proteins of this invention. Commensal bacteria which generate such hybrid proteins may be used to colonize an allergic patient. The result will be a constant

PCT/US93/02355

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source of allergens which will elicit the same protective immuned response as achieved with desensitization therapy.

The formation and expression of the hybrid protein M6:allergen 5 has been described above. Figs 11 and 12 show the results of immunization studies utilizing this hybrid surface antigen.

Fig. 11 illustrates the serum IgG response of mouse sera to purified allergen-5 in the hybrid surface protein. Mice were immunized either intradermally with <u>S. gordonii</u> expressing hybrid surface allergen-5 (10⁷ CFU) in Freund's adjuvant or intranasally with the same organisms. Animals were bled at 4, 6, 8, and 10 weeks after immunization and the serum checked by ELISA for IgG to purified allergen-5. Both mice immunized intradermally (V1-V4, hatched bars) and animals immunized intranasally (V5-V8, closed bars) expressed antibodies to the allergen. Control animals immunized with gordonii without surface exposed allergen-5 showed no response to the allergen. All animals immunized intranasally with the recombinant <u>gordonii</u> remained colonized for at least

PCT/US93/02355

12 weeks as determined by swabs and still expressed the allergen 5 for this period. All assays were performed with a 1:50 dilution of sera.

Fig. 12 shows the IgA response to allergin-5 in intranasal and intradermally immunized animals. Twelve weeks after immunization (intradermally or intranasally) with gordonii expressing allergin-5 (see Fig. 11) saliva was taken and the animals sacrificed to obtain lung and gut washes. The saliva and washes were analyzed for the presence of IgA specific for allergen-5 by ELISA. Results revealed that salivary IgA was present in the intradermal and intranasal immunized animals with a better response in the intranasally colonized animals. The best IgA response was seen in the lung washes of the intranasally colonized animals. While an IgA response was seen in the small intestine of the gut (S.I.), they were lower than the lung washes. All assays were performed with a 1:1 dilution of the samples.

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A particular advantage of the systems for delivery of the hybrid surface proteins of this inventions is that since the transformed organisms are normal commensals, they will continue to colonize and

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act as a constant stimulus of a protective immune response thereby avoiding the need of continual immunization.

The skilled artisan can readily conceive of numerous other applications of the novel products of this invention. For example, non-pathogenic gram positive bacteria can be produced to deliver surface antigens which will elicit protective antibodies against a wide variety of disease states including malaria, measles, pertussis and retroviral infections such as AIDS, coccidiosis, distemper, fowl pox and bruccelosis.

Another particular advantage of the delivery system of this invention when used as vaccine is that live bacteria are employed. As is known, live bacteria stimulate a much stronger immune response than attenuated or dead bacteria. Still another advantage is that, because non-pathogenic bacteria are employed, the vaccine is safe. Still another is that several methods of administration are possible.

- 42 -

The invention is not limited to <u>S. gordonii</u> as the carrier bacteria or even to streptococcus, in fact other non-pathogenic gram-positive bacteria may often be preferred. For example, bacteria normally present on the intestinal mucosa (<u>Enterococcus fecalis</u>) may be preferred to deliver an active polypeptide to the intestine.

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Moreover, the invention is not limited to the surface expression of one hybrid surface protein containing only one epitope. The carrier organism may be transformed by well known procedures to express a plurality of hybrid surface proteins each with a different active polypeptide or, alternatively, to express a hybrid surface protein in which the active polypeptide includes more than one epitope.

Thus far this invention has been described and illustrated principally with respect to the delivery of antigens to elicit protective antibodies but, as indicated above, the invention is not so limited. For example the invention can be used to deliver enzymes.

Humans deficient in the enzyme lactase which is secreted in the small intestine are unable to properly hydrolyze lactose into its components, D-glucose and D-galactose. One result is that they are unable to digest milk or milk products properly. Heretofore, the treatment has been life long ingestion of tablets or other therapeutic dosage forms containing the missing enzyme. In accordance with the practice of this invention, commensal intestinal bacteria are provided which are cloned to produce a hybrid protein, the active polypeptide of which is lactase. The bacteria may be employed to colonize the intestinal mucosa for the continual production of the required enzyme.

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Streptococcus mutans are major organisms responsible for tooth decay. They secrete glucosyltransferases which hydrolyze sucrose to produce glucose. The glucose, in turn, is formed into glucan polymers which adhere to the tooth surfaces and participate in the decay process. Utilizing the procedures described and claimed herein, S. mutans can be transformed to express hybrid surface proteins containing glucanase enzymes. The transformed bacteria may be employed to colonize the oral cavity

PCT/US93/02355

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and inhibit the formation of glucan polymers thereby protecting against tooth decay. Alternatively, Streptococcus mutans may be transformed to produce enzymes that digest dental plaque, thus preventing the initiation of decay.

The process of this invention substantially as describes above was employed utilizing Staphyloccus aureus as the bacteria for expression of the surface protein. When the anchor region (containing the LPXTGX segment, hydrophobic segment and charged tail segment) of Protein A, a surface molecule of this bacteria was fused to the enzyme alkaline phosphatase (a dimeric periplasmic protein of E. Coli) and the gene for this fusion placed back in S. aureus, the molecule was found to be translocated to the outside of the cell and anchored to its surface. The details of this procedured are found in Reference 65 which is incorporated herein by reference.

The skilled artisan will recognize that several variations of this invention are possible without departing from its spirit or scope.

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One variation which is particularly useful, especially for the production of vaccines is to include an adjuvant in the hybrid protein and the necessary nucleotide sequence for the expression of the desired protein in the gene used to generate the protein. Typically, a plasmid could be produced in which the gene for expressing the cholera toxin B subunit which is a known mucosal adjuvant is inserted between the C-region polypeptide and the active polypeptide. Alternatively, the gene for the subunit could be inserted upstream of the active polypeptide. This orientation may elicit an enhanced IgA response because in the usual interaction of CTB and the cell which produces antibodies, the CTB directly contacts the cell and stimulates the production of antibodies for the antigen for which CTB is functioning as an adjuvant. The plasmid may be employed in the double cross-over technique illustrated in Figs. 4 and 5 to incorporate the hybrid gene into the chromosome of a gram positive bacteria for the expression of a hybrid protein of the invention on the bacterial surface. Other immune enhancing proteins may be used instead of the cholera toxin B subunit to enhance the immune response to an active polypeptide.

The transformed bacteria of this invention are useful in diagnostic tests. They may be used, for example, to test for the presence of antibodies characteristic of a specific infection in the plasma of infected animals.

Conventionally, the procedure of testing for an infection caused by, for example, Neisseria gonorrhoeae requires the isolation and purification of an antigen which will react with a specific antibody characteristic of the infecting microorgansim and suspected to be present in the plasma. The antigen is then incubated with plasma or other body fluid. If a reaction takes place, it can be recognized by any of a variety of known tests.

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In the standard enzyme linked immunoassay procedure, the isolated and purified antigen is adhered to a solid phase glass or plastic substrate. It is then exposed to the body fluid such as plasma or serum. If the fluid contains an antibody characteristic of the infecting organism, an antigen/antibody reaction will take place. The

reaction product will remain on the solid phase which is then incubated with an anti-antibody tagged with a detectable enzyme such as horseradish peroxidase.

The transformed bacteria of this invention can be employed as the source of the antigen used in the diagnostic test. In such tests, the transformed bacteria is employed as an antigen. The skilled artisan will be familiar with several diagnostic tests of this nature. Such use, in accordance with this invention, eliminates the necessity of isolating and purifying the antigen as required by prior art methods.

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For diagnostic testing, a transformed bacteria of the invention is prepared in which the active polypeptide includes an epitope for a specific antibody which is characteristic of the microorganism causing the infection. The bacteria is adhered to the solid phase and the balance of the diagnostic test is conducted in the usual manner. Of course the test is not limited to the use of enzymes to detect the presence of antibody utilizing, for example, the ELISA assay. Other labels such a radioactive isotopes can be employed.

- 48 -

Diagnostic kits can be provided containing the transformed bacteria, typically in a lyophilized form to be reconstituted, together with the other reagents normally provided with such kits. These may include an aqueous buffer, a labeled anti-antibody, sterilized water and, possibly, other reagents. Alternatively, the transformed bacteria, either in aqueous suspension of lyophilized may be provided for use in the diagnostic test together with other reagents to be provided by the laboratory conducting the diagnostic test.

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The various segments of the hybrid surface proteins of the invention are not necessarily adjacent to each other in the protein or in the gene used to produce it. It may be convenient for any of a number of reasons apparent to the skilled artisan to separate the nucleotides on the hybrid gene used to express the protein by insertion of other nucleotides thereby to express hybrid proteins in which the active polypeptide and the cell associated polypeptide are separated by a selected spacer. It is not essential that the complete segment from the cell associated region of the surface antigen be employed in constructing the hybrid surface protein of the

invention so long as an anchor region beginning with the LPXTGX segment and ending at the charged tail segment is present to anchor the hybrid polypeptide to the bacteria that expresses it.

Fig. 9 shows one especially useful embodiment of this invention. It illustrates a universal plasmid containing a polylinker site. A polylinker is an oligonucleotide or DNA sequence which contains several restriction sites and therefore can be employed for the insertion of genes excised by a variety of restriction enzymes. For example, the plasmid can be constructed with a polylinker insert containing several restriction sites such as EcoRI, ClaI, HindIII, XboI, PstI, BamHI or any of several others.

Such polylinkers are either commercially available or may be readily constructed by the artisan.

To produce a plasmid of this invention the polylinker insert will be ligated into a plasmid already containing the selected gene segment for expressing the desired cell associated segment of the hybrid protein to be produced. The plasmid is shown as containing the leader sequence for the hybrid antigen, but such sequence is not necessarily in the

plasmid. It may be included in the oligonucleotide sequence used to express the active polypeptide.

Although the various gene segments in the plasmid are shown as adjacent to each other, they may be separated by spacers.

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The plasmid is shown as containing two different antibiotic resistant regions to assist in selection.

One, which is located within the polylinker site, separates it into two sections and will be removed when replaced by an insert. Other markers such as those resulting in color may be employed. Other formulations of a plasmid with polylinker sites may be constructed as required.

The plasmid without the marker which separates the linker, but with an effective promoter, can be employed to transform a gram positive bacteria. The plasmid shown can cross over with a chromosome as described above. A transformed bacteria containing the plasmid or the chromosome will express the desired surface hybrid protein.

- 51 -

The plasmid M6.1 367-441 which is described and claimed in the above identified parent application is especially useful in the practice of this invention. The preparation of this plasmid is described in the parent application. It has been deposited in <u>E. colistrain K561</u> at the American Type Culture Collection under the accession number ATCC68235.

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As explained in the parent applications, this plasmid can be fused with an appropriate DNA sequence and the hybrid gene inserted in an appropriate carrier, such as E. colic or a selected non-pathogenic gram positive bacteria. The resulting transformed bacteria will express a polypeptide of this invention. The transformed gram positive bacteria may be used directly as a vaccine. Alternatively, it can be used in a cross-over reaction as explained above to produce another transformed bacteria which will express the desired polypeptide more efficiently. The plasmid may also be used for the production of a unviersal peptide by the insertion of a polylinker.

A special advantage of the invention is that the recombinant gram positive bacteria may be delivered to the animal in need at the mucosal site at which the

- 52 -

pathogen normally invades the body. These includes oral as well as nasal, intestinal and vaginal sites. Such delivery will stimulate a local immune response at the site and be an effective means to raise a protective immune response to the pathogen. Since the bacteria employed will be non-pathogenic, normal organisms for that particular site, they may be employed without danger of toxic effects. The selected bacteria can be administered to fish or wild animals by mixing it in the food they eat or the water they inhabit.

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Utilizing the same procedures described above, hybrid proteins can be prepared containing the antibody eliciting antigens shown in the following Table 2 as active polypeptides. The table shows the polypeptide, which will produce antibodies protective against infection by a pathogen, the disease to be protected against and the publication which describes the polypeptide. In some instances the gene used to elicit the oligonucleotide coding for a peptide will need to be prepared and inserted in a plasmid. The procedures are completely parallel to the procedures described above. The C-terminal anchor region can be any of those specifically suggested above, or any

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other anchoring segment of a surface protein from a gram-positive bacteria described in Table 1. The active polypeptide of the transformed bacteria will include the peptide shown, fused to the leader sequence and a small segment of the N-terminus of a surface protein to permit proper translocation to the cell surface.

TABLE 2

PEPTIDE SEQUENCES SUITABLE FOR DEVELOPMENT OF VACCINES FROM HYBRID ANTIGENS

Peptide Disease (protein) Ref

A. H-(ASN-Ala-ASN-Pro)n-OH n 3 Malaria, cs protein 39
of Plasmodium falciparum

15 B. H-(Gly-Asp-Arg-Ala-Asp-GlyGln-Pro-Ala)n-OH n 2 of Plasmodium vivax

	c.	Glu-Gln-Asn-Val-Glu-His-	Malaria, Pf 155 of 41
		Asp-Ala	Plasmodium falciparum
			Y ·
	D.	Asn-Ala-Glu-Asn-Lys-Glu-	Malaria, Merozoite surface 42
		Glu-Leu-Thr-Ser-Ser-Asp-	protein of Plasmodium
5		Pro-Glu-Gly-Gin-Ile-Met	falciparum
	E.	Met-Gln-Trp-Asr-Ser-Thr-	Hepatitis, pre S(1) 43
	•	Ala-Phe-His-Gln-Thr-Leu-	
		Gln-Asp-Pro-Arg-Val-Arg-	•
		Gly-Leu-Tyr-Leu-Tyr-Leu-	
10		Pro-Ala-Gly-Gly	
	F .	Asp-Pro-Arg-Val-Arg-Gly-	Hepatitis, pre S(2) 44
		Leu-Tyr-Phe-Pro-Ala-Gly-	
		Gly-Ser-Ser-Gly-Thr-Val	
	G.	Cys-Thr-Lys-Pro-Thr-Asp-	Hepatitis Surface 45,46
15		Gly-Asn-Cys-Thr-Cys	antigen
	H.	Tyr-Ser-Thr-Leu-Tyr-Arg-	Poliovirus, replicase 47
		Trp-Leu-Asp-Asp-Ser-Phe	protein
			_

	Ι.	Asn-Ata-Pro-Ser-Lys-Inr-	Poliovirus, replicase	48
		Lys-Leu-Glu-Pro-Ser-Ala-	protein	
*		Phe		
,			l	
	J.	Lys-Lys-Pro-Asn-Val-Pro-	Poliovirus, VPg	49
5 .	•	Thr-Ile-Arg-Thr-Ala-Lys-		
		Val-Gln	- -	
	K.	Gly-Ser-Gly-Val-Arg-Gly-	Foot and Mouth Disease	50
		Asp-Ser-Gly-Ser-Leu-Ala-	VPI	
		Leu-Arg-Val-Ala-Arg-Gln-	•	
10		Leu-Pro		
	L.	Arg-His-Lys-Gln-Lys-Ile-	Foot and Mouth Disease	51
		Val-Ala-Pro-Val-Lys-Gln-	VPI	
		Thr-Leu		
		·		
	M.	Gly-Leu-Phe-Gly-Ala-Ile-	Influenza, HA2	52
15		Ala-Gly-Phe-Ile-Glu	Hemagglutinin protein	

- 56 -

The transformed bacteria of the invention are, in effect, therapeutic agents, and will be treated as such. They may be used alone or in association with pharmaceutically acceptable carriers of the type usually employed with such therapeutic agents. For oral or nasal administration, they may be suspended in aqueous isotonic saline which may be flavored or colored. For intestinal delivery, they may be placed in a capsule to deliver the transformed bacteria orally. Other methods of administration will be readily apparent to the skilled artisan and are within the scope of the invention.

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Since the products of the invention are utilized as live transformed bacteria to colonize the selected site of administration, no specific dosage is required. Typically, the selected dosage form will contain from about 10⁶ to 10¹⁰ organisms per dosage unit.

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Each of the publications cited is hereby incorporated by reference.

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WHAT IS CLAIMED IS:

- 1. A gene sequence which codes for a hybrid surface protein of a gram positive bacteria, said protein being attached to the carboxy terminus of the anchor region of a surface antigen normally expressed by the bacteria, the balance of the hybrid surface protein including an active polypeptide which may be delivered to an animal host for a useful purpose.
- 2. A plasmid comprising the gene sequence claimed in claim 1.
- 3. A chromosome comprising the gene sequence claimed in claim 1.
 - 4. The plasmid pVMB20.
 - 5. The plasmid pVMB21.
 - 6. The chromosome GP246.
 - 7. The plasmid M6.1.
- 8. A hybrid surface protein including the anchor sequence of a surface antigen normally expressed by a gram positive bacteria fused to an active polypeptide which may be delivered to a mammalian host for a useful purpose.
- 9. A hybrid surface protein of claim 8 wherein the active polypeptide is an enzyme.

- 10. A hybrid surface protien of claim 8 wherein the active polypeptide is a surface antigen of a mammalian tumor cell.
- 11. A hybrid surface protein of claim 8 wherein the active polypeptide is a surface antigen of male sperm.

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- 12. A hybrid surface prtein of claim 8 wherein the active polypeptide is an allergen.
- 13. A hybrid surface protein of claim 8 wherein the active polypeptide includes an antigenic determinant of a surface antigen of a bacteria, virus, parasite or fungus.
- 14. A hybrid surface protein of claim 8 wherein the anchor sequence is the anchor sequence of a streptococcal M protein.
- 15. A hybrid surface protein claim 14 wherein the active polypeptide includes an antigenic determinant of a surface antigen of a bacteria, virus, parasite or fungus.
 - 16. The hybrid surface protein M6:E7.
 - 17. The hybrid surface protein M6:allergen 5.
- 18. A non-pathogenic gram positive bacteria which expresses a hybrid surface protein having an anchor sequence which is attached to the bacteria and is the anchor sequence of a surface antigen normally

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expressed by a gram positive bacteria fused to an active polypeptide which may be delivered to an animal host for a useful purpose.

- 19. A non-pathogenic gram-positive bacteria of claim 18 wherein the active polypeptide is an enzyme.
- 20. A non-pathogenic gram-positive bacteria of claim 18 wherein the active polypeptide is a surface antigen of a mammalian tumor cell.
- 21. A non-pathogenic gram-positive bacteria of claim 18 wherein the active polypeptide is a surface antigen of male sperm.
- 22. A non-pathogenic gram-positive bacteria of claim 18 wherein the active polypeptide is an allergen.
- 23. A non-pathogenic gram-positive bacteria of claim 18 wherein the active polypeptide includes an antigenic determinant of a surface antigen of a bacteria, virus, parasite or fungus.
 - 24. A non-pathogenic gram-positive bacteria of claim 18 wherein the anchor sequence is the anchor sequence of a streptococcal M protein.
 - 25. A non-pathogenic gram-positive bacteria of claim 24 wherein the active polypeptide includes an antigenic determinant of a surface antigen of a bacteria, virus, parasite or fungus.

- 26. A non-pathogenic gram-positive bacteria of claim 18 wherein the hybrid surface protein is expressed by a plasmid.
- 27. A non-pathogenic gram-positive bacteria of claim 18 wherein the hybrid surface protein is expressed by a chromosomal gene.
- 28. <u>Streptococcus gordonii</u> GP246 which expresses the hybrid antigen M6:E7.
- 29. <u>Streptococcus gordonii</u> GP246 which expresses the hybrid antigen M6:E7 utilizing the plasmid pVMB21.
- 30. <u>Streptococcus gordonii</u> GP246 which expresses the hybrid antigen M6:E7 utilizing the chromosone GP246.
- 31. A vaccine to protect an animal host against infection by a pathogenic bacteria which comprises a pharmaceutically acceptable carrier and a non-pathogenic gram positive bacteria which expresses a hybrid surface protein, said protein being attached to the bacteria at the carboxy terminus of the anchor sequence of a surface antigen normally expressed by a gram-positive bacteria, the balance of the hybrid surface protein being an active polypeptide which includes the antigenic determinant of a surface antigen of a pathogenic bacteria to stimulate the

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production of protective antibodies in an amount sufficient to inhibit infection of an animal host by said pathogenic bacteria.

- 32. A vaccine of claim 31 wherein the active polypeptide includes the antigenic determinant of a surface antigen of a streptococcus.
- 33. A vaccine of claim 31 wherein the non-pathogenic bacteria is Streptococcus gordonii.
- 34. A vaccine of claim 31 wherein the non-pathogenic bacteria is <u>Streptococcus gordonii</u> transformed to express the hybrid surface protein M6:E7.
- 35. A vaccine of claim 31 wherein the Streptococcus is transformed by the plasmid pVMB21.
- 36. A vaccine of claim 31 wherein the Streptococcus gordonii is transformed by the chromosone GP246.
- 37. A method of protecting against infections by a gram-positive pathogenic bacteria in an animal in need of such protection which comprises administration of a vaccine of claim 31.

- 70 -

38. A plasmid comprising a gene segment for expressing the anchor sequence of a surface antigen normally expressed by a gram positive bacteria together with a polylinker segment which is separated into two sections by a marker.

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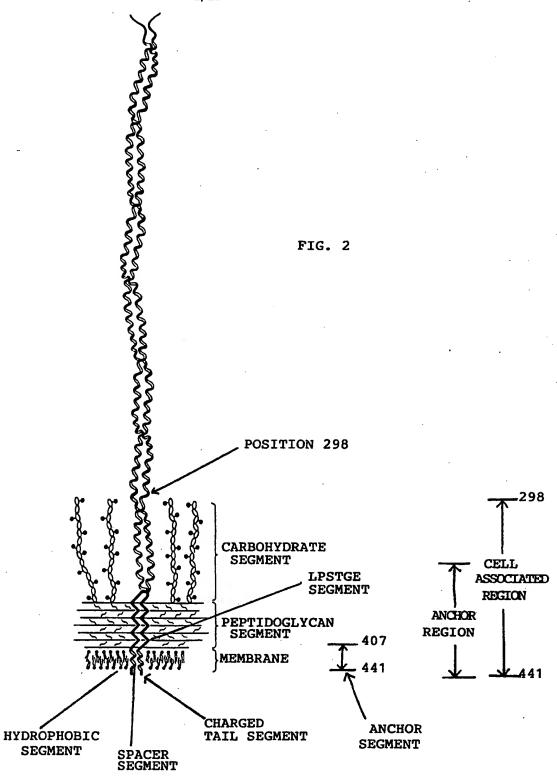
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- 39. A method of testing for an infection which is characterized by the presence of antibodies in a body fluid of an animal, said antibodies being characteristic of the infecting organism, said test comprising incubating the body fluid with a non-pathogenic gram positive bacteria which expresses a hybrid surface protein the anchor sequence of which is attached to the bacteria and is the anchor sequence of a surface antigen normally expressed by gram positive bacteria, the balance of the hybrid surface protein including an antigen which will react with said antibodies and thereafter, determining if a reaction has taken place.
- 40. A kit useful for testing to determine the presence of an infection, said infection being characterized by the presence of antibodies in a body fluid of an animal, said antibodies being characteristic of the infecting organism, said kit containing a non-pathogenic gram positive bacteria which expresses a hybrid surface protein the anchor

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sequence of which is attached to the bacteria and is the anchor sequence of a surface antigen normally expressed by gram positive bacteria, the balance of the hybrid surface protein including an antigen which will react with said antibodies.

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I . Arg Val Phe Pro Arg Gly Thr
                          Val Glu Asn l'en
                          Asp Lys Ala Arg Clu Leu
                                  c de
                         Leu Asn Lys Tyr Asp Val Glu
                         Ann Ser Met Leu Cin Ala Ann
                    32
                         Asn Asp Lys Leu Thr Thr Glu
                    39
                         Asn Asn Asn Leu The Asp Gln
                         Asa Lye Asa Leu The The Glu
                    53
                         Asn Lys Asn Leu The Asp Gin
                         Asn Lys Asn Leu Thr Thr Glu
                    60
                         Asn Lys Asn Leu Thr Asp Gln
                    67
                    74
                         Asn Lys Asn Leu The The Glu
                    81
                         Asn Lys Glu Leu Lys Ala Glu
                         Glu Asn Arg Leu Thr Thr Glu
                         Asn Lys Gly Leu Thr Lys Lys
                    95
                   102
                         Les Ser Glu Ala Glu Glu Glu
                   109
                         Ala
                         Ala Asn. Lys Glu Arg Glu Asn
                   110
                         Lys Glu Ala Ile Gly Thr Leu
                   117
                         Lys Lys Thr Leu Asp Glu Thr
                   124
                                     Val Lys Asp Lys
                   131
                   135
                         lle Ala Lys Glu Gin Glu Ser
                   142
                         Lys Clu Thr lie Cly Thr Leu
                   149
                         Lys Lys Thr Leu Asp Glu Thr
                   156
                                     Val Lys Asp Lys
                         ile Ala Lys Glu Gin Glu Ser
                         Lys Glu Thr lie Gly The Leu
                   167
                                                                             FIG. 3
                         Lys Lys Thr Leu Asp Glo Thr
                   174
                  181
                                     Val Lys Asp Lys
                         lle Ala Lys Glu Gln Glu Ser
                  185
                        Lys Glu Thr Ile Gly Thr Leu
Lys Lys Ile Leu Asp Glu Thr
                  192
                  199
                                     Val Lys Asp Lys
                  206
                  210
                         lle Ala Arg Glu Gla Lys Ser
                  217
                        Lys Gin Asp IIe Gly Als Lee
                  224
                         Lys Gin Giu Leu Ala Lys Lys
                  231
                                         Asp Clu Cly
                  234
                        Ass Lys Val Ser Clu Als Ser
                        Arg Lys Gly Leu Arg Arg Asp
                  241
                  248
                        Leu Asp Ala Ser Arg Glu Ala
                        Lys Lys Gin Val Glu Lys Asp
                  255
                        Leu Ala Asn Leu The Ala Glu
                  282
                        Les Asp Lys Val Lys Glu Glu
                  269
                  275
                        Lys Gla lle Ser Asp Ala Ser
                  283
                        Arg Gla Gly Leu Arg Arg Asp
                  290
                        Leu Asp Ala Ser Arg Glu Ala
                        Lys Lys Gin Val Glu Lys Ala
Leu Glu Glu Ala Asn Ser Lys
POSITION 298
                 207
                                                                                          CARBOHYDRATE
                  304
                                                                                             SEGMENT
                        Les Ala Ala Les Glo Lys Les
                  311
                  318
                        Asa Lys Glu Leu Glu Glu Ser
                  325
                        Lys Lys Leu Thr Glu Lys Glu
                  332
                        Lys Ala Glu Leu Gla
                  337
                           Ala Lys Leu Glu Ala Glu
                                                                              CELL-ASSOCIATED
                 343
                        Ala Lys Ala Leu Lys Glu Gin
                        Les Ala Lys Cln Ala Glu Clu
                 350
                                                                              REGION
                 357
                        Lee Ala Lys Leu Arg Ala
                 363
                        Gly Lys Ala Ser Asp Ser Gin Thr
                 371
                        Pro Asp Ala Lya
                 375
                        Pro Gly Asn Lys Val Val
                 381
                        Pro Gly Lys Gly Gla Ala
                                                                            PEPTIOGLYCAN SEGMENT
                 387
                        Pro Gin Ala Gly The Lys
                       Pro Asn Gin Asn Lys Ala
Pro Met Lys Giu Thr Lys Arg Gin Leu
Pro Ser Thr Gly Giu Thr Ala Asn Pro
                 303
                 398
    LPSTGE
                 408
                       Phe Phe Thr Ala Ala Ala Leu
Thr Val Met Ala Thr Ala Gly
Val Ala Ala Val Val

HYDROPHOBIC
SEGMENT
                 417
                                                                               ANCHOR REGION
                 424
                 431
                 438
                       Lys Arg Lys Glu Clu Asn
                                                                CHARGED TAIL SEGMENT
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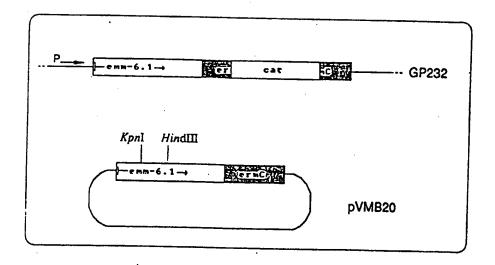


Fig. 4

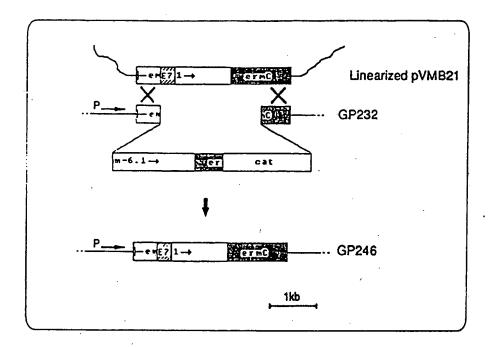


Fig. 5

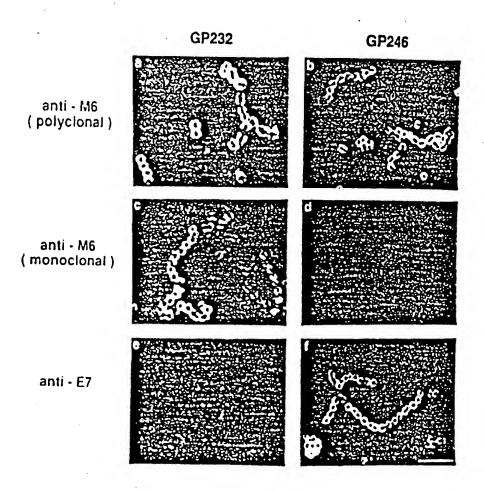


FIG. 6

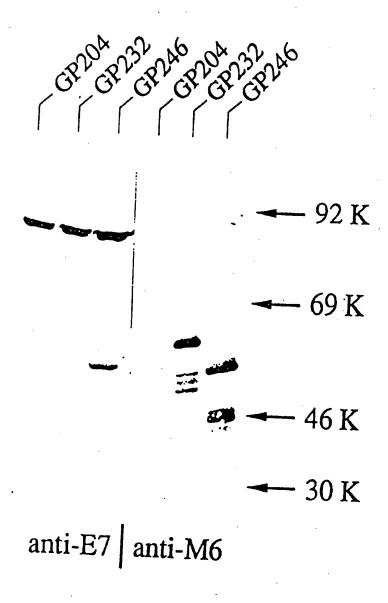


Fig. 7

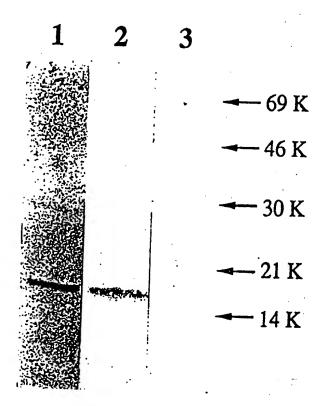


Fig. 8

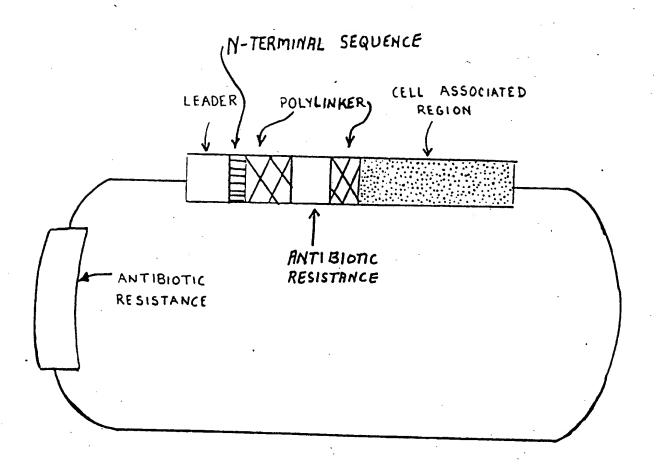


Fig. 9

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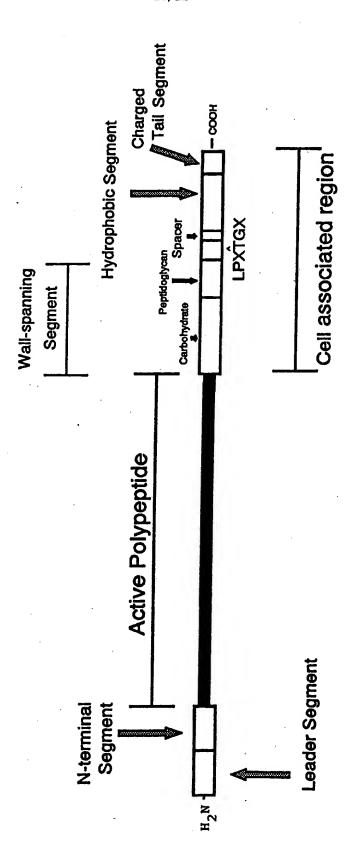
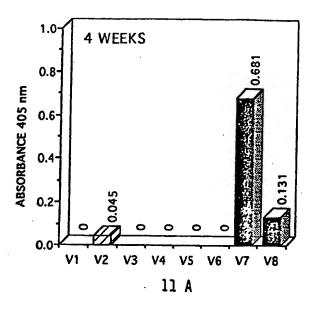
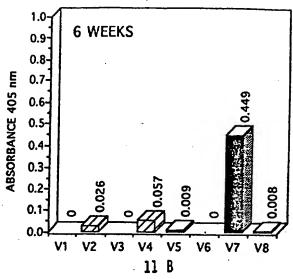
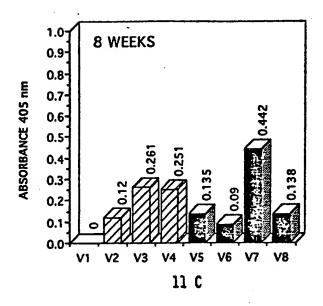
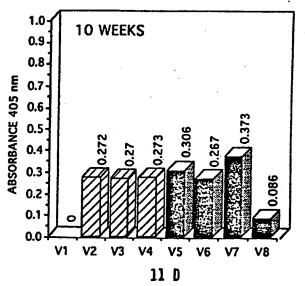


FIG. 10

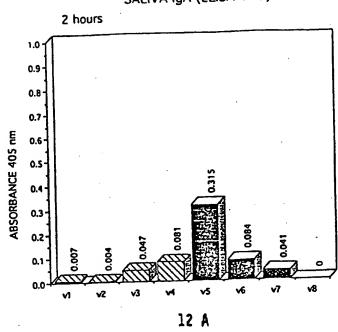


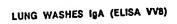






SALIVA IgA (ELISA VV8)





2 hours

GUT WASHES S.I. (ELISA VV8)

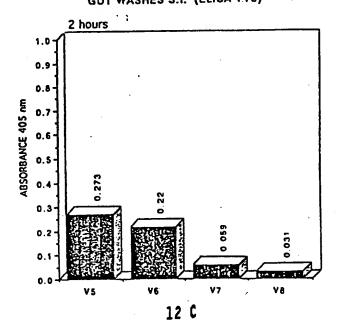


FIGURE 12 A - C

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